



RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number: 10/004,115
Source: OIPE
Date Processed by STIC: 8/12/0

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- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
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Revised 01/29/2002



OIPE

100% Reply
100%RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/004,115DATE: 08/12/2002
TIME: 16:10:55Input Set : A:\72249.txt
Output Set: N:\CRF4\08122002\J004115.raw

4 <110> APPLICANT: ASAKO, HIROYUKI
 5 MATSUMURA, KENJI
 6 SHIMIZU, MASATOSHI
 7 ITO, NOBUYA
 8 WAKITA, RYUHEI
 10 <120> TITLE OF INVENTION: PROCESS FOR PRODUCING OPTICALLY ACTIVE
 11 4-HALO-3-HYDROXYBUTANOATE
 14 <130> FILE REFERENCE: 72249
 16 <140> CURRENT APPLICATION NUMBER: US 10/004,115
 C--> 17 <141> CURRENT FILING DATE: 2002-08-05
 19 <150> PRIOR APPLICATION NUMBER: JP 2000-372704
 20 <151> PRIOR FILING DATE: 2000-12-07
 22 <150> PRIOR APPLICATION NUMBER: JP 2001-006144
 23 <151> PRIOR FILING DATE: 2001-01-15
 25 <150> PRIOR APPLICATION NUMBER: JP 2001-026594
 26 <151> PRIOR FILING DATE: 2001-02-02
 28 <150> PRIOR APPLICATION NUMBER: JP 2001-175175
 29 <151> PRIOR FILING DATE: 2001-06-11
 31 <160> NUMBER OF SEQ ID NOS: 33
 33 <170> SOFTWARE: FastSEQ for Windows Version 3.0

ERRORED SEQUENCES

203 <210> SEQ ID NO: 6
 204 <211> LENGTH: 14
 205 <212> TYPE: PRT
 206 <213> ORGANISM: PENICILLIUM CITRINUM
 208 <400> SEQUENCE: 6
 E--> 209 Tyr Glu Asp Val Leu Xaa Xaa Ile Asp Asp Ser Leu Lys Arg
 210 1 5 10
 E--> 212 <210> SEQ ID NO: 8 } Skipped sequence - step 3
 296 <210> SEQ ID NO: 15
 297 <211> LENGTH: 697
 298 <212> TYPE: DNA
 299 <213> ORGANISM: ESCHERICHIA COLI
 301 <400> SEQUENCE: 15
 E--> 302 cgctctaaaa ctantggatc ccccggtctg caggaattcg gcggcgccgg atccaaacgga 60
 E--> 303 aanactttca cactgagcaa cggcgtcaaa attcctggcg tcggctttgg tacctncgct 120
 E--> 304 agtgaaggtt ccaagggtcga aacctatnct gctgtcacca ctgcccgtaa aaccgggttac 180
 E--> 305 cgtcncttgg actgtgcctg gtactacctg aacaagggtg aggttggtaa ggtntccgt 240
 306 gacttcctga agggaaaaccc ctcgggtaaag cgtgaggaca tcttcgtctg caccaggta 300
 E--> 307 tggaaaccacc tccaccgtta tgaggacgtc ctctggtcca ttgacnactc cctgaagcgt 360

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308	cttggacttg actacgttga tatgttctc gttcaactggc ccattgtgc cgaaaaaaat	420
E--> 309	ggccagggtg agccaaaaat tggccctgac ggcaaaatacn tcnttctcaa ggacctgacc	480
E--> 310	gaaanccna ncccacctgg cgcgctatgg aaaaaatttn tgangatccc aaggccaggt	540
E--> 311	ccattggtgtt ttccaattgg accattgccc accttgagaa gatgtccaag ttnccaagg	600
E--> 312	tnatgcctca cgccaaccag atcgagattc accccttcct gccaacgag gagctggtgc	660
E--> 313	agtactgctt ttccaagaac antatgccc tagcgta	697
340	<210> SEQ ID NO: 18	
341	<211> LENGTH: 331	
342	<212> TYPE: DNA	
343	<213> ORGANISM: ESCHERICHIA COLI	
345	<400> SEQUENCE: 18	
E--> 346	cgctctaaaa ctantggatc ccccggtcg caggaattcg gggccgcgg atccttcata	60
347	cccatcatgt ctaacggaaa gactttaca ttgagcaacg gctcaagat tcctggcg	120
348	ggcttggta ccttcgttag tgaaggttcc aaggccgaga cctataactgc tgtcaccact	180
349	gcccgtgaaga ccgttaccg tcacttggac tgcctgtt actacgtaa cgagggtgag	240
350	gttggtgagg gtatccgtga cttcctgaag gagaaccctt cggtgaagcg tgaggacatc	300
351	ttcgtctgca ccaagggtgt gaaccaccc c	331
353	<210> SEQ ID NO: 19	
354	<211> LENGTH: 743	
355	<212> TYPE: DNA	
356	<213> ORGANISM: ESCHERICHIA COLI	
358	<400> SEQUENCE: 19	
359	caaccagatc gagattcacc cttcctgcc caacgaggag ctggtgcaatc actgcttc	60
360	caagaacatt atgcccgtgg cctactctcc tctggctcg cagaaccagg ttcccaccac	120
361	cggtgagcgg gtcagcgaga acaagactct gaacgagatc gcccggaaagg gcccggacac	180
362	ccttgctca gtttttattt cctgggtct gcccgtggc tacgtcttc tccccaaagag	240
363	ctccaacccc aagcgattt agtccaactt caagaggttcc gagctctccg atgcccactt	300
364	tgaagccatc aatgccgtt ccaagggtcg tcacttccgt ttctcaaca tgaaggatac	360
365	tttcggatata gatgtctggc ccgaggagac gccaagaac ctgtctcgat gatctctac	420
E--> 366	gaaattataa aatnacacccn acnaaaancc aaagcgaaag gatgtatccc aaaantttt	480
E--> 367	agggtttctt ggtgaaaac gtttattttt cccgaantga angaataat ganctngatt	540
E--> 368	tctccaaaaa aaaaaaaaaa aaaaacggtc cgccggccgtt ccnnnggggg gcccgggttcc	600
E--> 369	caattcnccc cttatnattt aattttttt taanggggnc aaattccncc nnattccnt	660
E--> 370	cnaatttgn nggccgcctc caaacttccn tcntnaaagg gncccaattc ccccccatt	720
E--> 371	aantggantt cctntttacc ttt	743
E--> 657	<210> SEQ ID NO: 32	

pls in sequence

Seq p. 4

{ Seq p. 3 - Skipped sequence

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/004,115

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Input Set : A:\72249.txt
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Skipped Sequences (NEW RULES):

Sequence(s) missing. If intentional, please use the following format for each skipped sequence.

<210> sequence id number

<400> sequence id number

000

Seq# : 7, 31

VARIABLE LOCATION SUMMARY
PATENT APPLICATION: US/10/004,115

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TIME: 16:10:56

Input Set : A:\72249.txt
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Use of n's or Xaa's(NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which
residue n or Xaa represents.

Seq#:6; Xaa Pos. 6,7
Seq#:9; N Pos. 3,6,9
Seq#:10; N Pos. 3,6,9
Seq#:11; N Pos. 3,6,9
Seq#:12; N Pos. 3,6,9
Seq#:13; N Pos. 3,6,9
Seq#:14; N Pos. 3,6,9
Seq#:15; N Pos. 14,63,116,148,185,235,346,460,463,485,489,491,520,524,593
Seq#:15; N Pos. 602,682
Seq#:18; N Pos. 14
Seq#:19; N Pos. 434,440,443,448,458,467,475,506,510,517,522,533,535,583,584
Seq#:19; N Pos. 607,616,634,639,648,651,652,659,662,664,670,671,690,693,695
Seq#:19; N Pos. 702,717,723,728,734

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/004,115

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Input Set : A:\72249.txt
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L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:39 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:201 M:300 W: (50) Intentionally skipped Sequence. : Sequence Id (5) SEQUENCE:
L:209 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:6
L:212 M:214 E: (33) Seq.# missing, SEQ ID NO:7
L:234 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:9
L:234 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:9
L:234 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:246 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:10
L:246 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:10
L:246 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:258 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:11
L:258 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:11
L:258 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:270 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:12
L:270 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:12
L:270 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:282 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:13
L:282 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
L:282 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
L:294 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:14
L:294 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:294 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:302 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:15
M:340 Repeated in SeqNo=15
L:346 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:18
L:366 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:19
M:340 Repeated in SeqNo=19
L:657 M:214 E: (33) Seq.# missing, SEQ ID NO:31